

Package: SLTCA (via r-universe)

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Type Package

Title SLTCA: Scalable and Robust Latent Trajectory Class Analysis
Using Artificial Likelihood

Description Conduct latent trajectory class analysis with longitudinal
observations.

Depends R (>= 3.3.0)

Imports stats, geepack, VGAM, Matrix

Version 0.1.0

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Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

Repository <https://tengfei-emory.r-universe.dev>

RemoteUrl <https://github.com/tengfei-emory/sltca>

RemoteRef HEAD

RemoteSha ba768bc73c8d8c57fb00d63a0da81c48a2057b01

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simulation	<i>Simulate a dataset which can be analyzed by SLTCA</i>
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Description

Simulate a dataset with longitudinal observations.

Usage

```
simulation(n)
```

Arguments

n	Sample size.
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Value

Returns a data frame with 6 longitudinal features y.1 - y.6, including count (y.1 and y.2), binary (y.3 and y.4) and continuous (y.5 and y.6) type. Variable baselinecov is the baseline risk factor of latent classes. Variable latent is the true latent class labels.

Author(s)

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Examples

```
dat <- simulation(500)
```

SLTCA	<i>SLTCA: Scalable and Robust Latent Trajectory Class Analysis Using Artificial Likelihood</i>
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Description

Conduct latent trajectory class analysis with longitudinal observations.

Usage

```
SLTCA(k = 20, dat, num_class, id, time, num_obs, features, Y_dist, covx, ipw,
stop, tol = 0.005, max = 50, varest = T, balanced = T, MSC = "EQIC",
verbose = T)
```

Arguments

<code>k</code>	Number of random initialization to start the algorithm.
<code>dat</code>	Input data matrix.
<code>num_class</code>	Number of latent classes in the fitted model.
<code>id</code>	Column name in the data matrix ‘dat’ for the patient id.
<code>time</code>	Column name in the data matrix ‘dat’ for the time of longitudinal observations.
<code>num_obs</code>	Column name in the data matrix ‘dat’ for the number of longitudinal observations (number of visits).
<code>features</code>	A vector of column names in the data matrix ‘dat’ for the longitudinal observations.
<code>Y_dist</code>	A vector indicating the type of longitudinal observations. An element of <code>Y_dist</code> can be ‘normal’, ‘bin’, and ‘poi’ for continuous, binary and count data.
<code>covx</code>	A vector of column names in the data matrix ‘dat’ for baseline latent class risk factors.
<code>ipw</code>	Column name in the data matrix ‘dat’ for the inverse probability weights for missingness. <code>ipw=1</code> if not specified.
<code>stop</code>	Stopping criterion for the algorithm. <code>stop</code> can be either ‘tau’ based on posterior probabilities or ‘par’ based on point estimation.
<code>tol</code>	A constant such that the algorithm stops if the stopping criterion is below this constant.
<code>max</code>	Maximum number of iterations if the algorithm does not converge.
<code>varest</code>	True or False: whether conduct variance estimation or not.
<code>balanced</code>	True or False: whether the longitudinal observations are equally spaced.
<code>MSC</code>	Model selection criteria: ‘AQIC’, ‘BQIC’ or ‘EQIC’.
<code>verbose</code>	Output progress of fitting the model.

Value

A list with point estimates (`alpha`, `beta0`, `beta1`, `phi`, `gamma`), variance estimates (ASE), posterior membership probabilities (`tau`) and QICs (`qic`) of the latent trajectory class model.

Author(s)

Teng Fei. Email: tfei@emory.edu

References

Hart, Fei and Hanfelt (202x), Scalable and Robust Latent Trajectory Class Analysis Using Artificial Likelihood (in press).

Examples

```
dat <- simulation(500)

res <- SLTCA(k=1,dat,num_class=2,"id","time","num_obs",paste("y.",1:6,sep=''),
  Y_dist=c('poi','poi','bin','bin','normal','normal'),
  "baselinecov",1,stop="tau",tol=0.005,max=50,
  varest=T,balanced=T,MSC='EQIC',verbose=T)
```

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